

Title: Inbred Maize Line PH48V

Anticipated Class: 800

Subclass: 200

Art Unit: 1649

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**Field of the Invention**

This invention is in the field of maize breeding, specifically relating to an inbred maize line designated PH48V.

**Background of the Invention**

The goal of plant breeding is to combine in a single variety or hybrid various desirable traits. For field crops, these traits may include resistance to diseases and insects, tolerance to heat and drought, reducing the time to crop maturity, greater yield, and better agronomic quality. With mechanical harvesting of many crops, uniformity of plant characteristics such as germination and stand establishment, growth rate, maturity, and plant and ear height, is important.

Field crops are bred through techniques that take advantage of the plant's method of pollination. A plant is self-pollinated if pollen from one flower is transferred to the same or another flower of the same plant. A plant is cross-pollinated if the pollen comes from a flower on a different plant.

Plants that have been self-pollinated and selected for type for many generations become homozygous at almost all gene loci and produce a uniform population of true breeding progeny. A cross between two different homozygous lines produces a uniform population of hybrid plants that may be heterozygous for many gene loci. A cross of two plants each heterozygous at a number of gene loci will produce a population of hybrid plants that differ genetically and will not be uniform.

Maize (*zea mays* L.), often referred to as corn in the United States, can be bred by both self-pollination and cross-pollination techniques. Maize has separate male and female flowers on the same plant, located on the tassel and the ear, respectively. Natural pollination occurs in maize when wind blows pollen from the tassels to the silks that protrude from the tops of the ears.

A reliable method of controlling male fertility in plants offers the opportunity for improved plant breeding. This is especially true for development of maize hybrids,

which relies upon some sort of male sterility system. There are several options for controlling male fertility available to breeders, such as: manual or mechanical emasculation (or detasseling), cytoplasmic male sterility, genetic male sterility, gametocides and the like.

5           Hybrid maize seed is typically produced by a male sterility system incorporating manual or mechanical detasseling. Alternate strips of two maize inbreds are planted in a field, and the pollen-bearing tassels are removed from one of the inbreds (female). Providing that there is sufficient isolation from sources of foreign maize pollen, the ears of the detasseled inbred will be fertilized only from the other inbred (male), and the  
10       resulting seed is therefore hybrid and will form hybrid plants.

          The laborious, and occasionally unreliable, detasseling process can be avoided by using cytoplasmic male-sterile (CMS) inbreds. Plants of a CMS inbred are male sterile as a result of factors resulting from the cytoplasmic, as opposed to the nuclear, genome. Thus, this characteristic is inherited exclusively through the female parent in  
15       maize plants, since only the female provides cytoplasm to the fertilized seed. CMS plants are fertilized with pollen from another inbred that is not male-sterile. Pollen from the second inbred may or may not contribute genes that make the hybrid plants male-fertile. Seed from detasseled fertile maize and CMS produced seed of the same hybrid can be blended to insure that adequate pollen loads are available for fertilization when  
20       the hybrid plants are grown.

          There are several methods of conferring genetic male sterility available, such as multiple mutant genes at separate locations within the genome that confer male sterility, as disclosed in U.S. Patents 4,654,465 and 4,727,219 to Brar et al. and chromosomal translocations as described by Patterson in U.S. Patents No. 3,861,709 and 3,710,511.  
25       These and all patents referred to are incorporated by reference. In addition to these methods, Albertsen et al., of Pioneer Hi-Bred, U.S. Patent No. 5,432,068, have developed a system of nuclear male sterility which includes: identifying a gene which is critical to male fertility; silencing this native gene which is critical to male fertility; removing the native promoter from the essential male fertility gene and replacing it with  
30       an inducible promoter; inserting this genetically engineered gene back into the plant; and thus creating a plant that is male sterile because the inducible promoter is not "on" resulting in the male fertility gene not being transcribed. Fertility is restored by inducing, or turning "on", the promoter, which in turn allows the gene that confers male fertility to be transcribed.

There are many other methods of conferring genetic male sterility in the art, each with its own benefits and drawbacks. These methods use a variety of approaches such as delivering into the plant a gene encoding a cytotoxic substance associated with a male tissue specific promoter or an antisense system in which a gene critical to fertility is identified and an antisense to that gene is inserted in the plant (see: Fabinjanski, et al. EPO 89/3010153.8 publication no. 329,308 and PCT application PCT/CA90/00037 published as WO 90/08828).

Another system useful in controlling male sterility makes use of gametocides. Gametocides are not a genetic system, but rather a topical application of chemicals. These chemicals affect cells that are critical to male fertility. The application of these chemicals affects fertility in the plants only for the growing season in which the gametocide is applied (see Carlson, Glenn R., U.S. Patent Number: 4,936,904). Application of the gametocide, timing of the application and genotype specificity often limit the usefulness of the approach.

#### **Development of Maize Inbred Lines**

The use of male sterile inbreds is but one factor in the production of maize hybrids. Plant breeding techniques known in the art and used in a maize plant breeding program include, but are not limited to, recurrent selection, backcrossing, pedigree breeding, restriction length polymorphism enhanced selection, genetic marker enhanced selection and transformation. The development of maize hybrids in a maize plant breeding program requires, in general, the development of homozygous inbred lines, the crossing of these lines, and the evaluation of the crosses. Pedigree breeding and recurrent selection breeding methods are used to develop inbred lines from breeding populations. Maize plant breeding programs combine the genetic backgrounds from two or more inbred lines or various other germplasm sources into breeding pools from which new inbred lines are developed by selfing and selection of desired phenotypes. The new inbreds are crossed with other inbred lines and the hybrids from these crosses are evaluated to determine which of those have commercial potential. Plant breeding and hybrid development, as practiced in a maize plant breeding program, are expensive and time consuming processes.

Pedigree breeding starts with the crossing of two genotypes, each of which may have one or more desirable characteristics that is lacking in the other or which complements the other. If the two original parents do not provide all the desired characteristics, other sources can be included in the breeding population. In the

pedigree method, superior plants are selfed and selected in successive generations. In the succeeding generations the heterozygous condition gives way to homogeneous lines as a result of self-pollination and selection. Typically in the pedigree method of breeding five or more generations of selfing and selection is practiced:  $F_1 \rightarrow F_2$ ;  $F_2 \rightarrow F_3$ ;

5  $F_3 \rightarrow F_4$ ;  $F_4 \rightarrow F_5$ , etc.

Recurrent selection breeding, backcrossing for example, can be used to improve an inbred line and a hybrid which is made using those inbreds. Backcrossing can be used to transfer a specific desirable trait from one inbred or source to an inbred that lacks that trait. This can be accomplished, for example, by first crossing a superior  
10 inbred (recurrent parent) to a donor inbred (non-recurrent parent), that carries the appropriate gene(s) for the trait in question. The progeny of this cross is then mated back to the superior recurrent parent followed by selection in the resultant progeny for the desired trait to be transferred from the non-recurrent parent. After five or more backcross generations with selection for the desired trait, the progeny will be  
15 homozygous for loci controlling the characteristic being transferred, but will be like the superior parent for essentially all other genes. The last backcross generation is then selfed to give pure breeding progeny for the gene(s) being transferred. A hybrid developed from inbreds containing the transferred gene(s) is essentially the same as a hybrid developed from the same inbreds without the transferred gene(s).

20 Elite inbred lines, that is, pure breeding, homozygous inbred lines, can also be used as starting materials for breeding or source populations from which to develop other inbred lines. These inbred lines derived from elite inbred lines can be developed using the pedigree breeding and recurrent selection breeding methods described earlier. As an example, when backcross breeding is used to create these derived lines in a  
25 maize plant breeding program, elite inbreds can be used as a parental line or starting material or source population and can serve as either the donor or recurrent parent.

### **Development of Maize Hybrids**

A single cross maize hybrid results from the cross of two inbred lines, each of which has a genotype that complements the genotype of the other. The hybrid progeny  
30 of the first generation is designated  $F_1$ . In the development of commercial hybrids in a maize plant breeding program, only the  $F_1$  hybrid plants are sought. Preferred  $F_1$  hybrids are more vigorous than their inbred parents. This hybrid vigor, or heterosis, can be manifested in many polygenic traits, including increased vegetative growth and increased yield.





Seed Science and Technology 14, pp. 1-8 (1995), the disclosure of which is expressly incorporated herein by reference. Through these technologies, the homozygosity of the self pollinated line can be verified by analyzing allelic composition at various loci along the genome. Those methods allow for rapid identification of the invention disclosed  
5 herein. See also, "Identification of Atypical Plants in Hybrid Maize Seed by Postcontrol and Electrophoresis" Sarca, V. et al., Probleme de Genetica Teoritica si Aplicata Vol. 20 (1) p. 29-42.

As is readily apparent to one skilled in the art, the foregoing are only some of the various ways by which the inbred can be obtained by those looking to use the  
10 germplasm. Other means are available, and the above examples are illustrative only.

Maize is an important and valuable field crop. Thus, a continuing goal of plant breeders is to develop high-yielding maize hybrids that are agronomically sound based on stable inbred lines. The reasons for this goal are obvious: to maximize the amount of grain produced with the inputs used and minimize susceptibility of the crop to pests  
15 and environmental stresses. To accomplish this goal, the maize breeder must select and develop superior inbred parental lines for producing hybrids. This requires identification and selection of genetically unique individuals that occur in a segregating population. The segregating population is the result of a combination of crossover events plus the independent assortment of specific combinations of alleles at many  
20 gene loci that results in specific genotypes. The probability of selecting any one individual with a specific genotype from a breeding cross is infinitesimal due to the large number of segregating genes and the unlimited recombinations of these genes, some of which may be closely linked. However, the genetic variation among individual progeny of a breeding cross allows for the identification of rare and valuable new genotypes.  
25 These new genotypes are neither predictable nor incremental in value, but rather the result of manifested genetic variation combined with selection methods, environments and the actions of the breeder.

Thus, even if the entire genotypes of the parents of the breeding cross were characterized and a desired genotype known, only a few if any individuals having the  
30 desired genotype may be found in a large segregating  $F_2$  population. Typically, however, neither the genotypes of the breeding cross parents nor the desired genotype to be selected is known in any detail. In addition to the preceding problem, it is not known how the genotype would react with the environment. This genotype by environment interaction is an important, yet unpredictable, factor in plant breeding. A

breeder of ordinary skill in the art cannot predict the genotype, how that genotype will interact with various climatic conditions or the resulting phenotypes of the developing lines, except perhaps in a very broad and general fashion. A breeder of ordinary skill in the art would also be unable to recreate the same line twice from the very same original  
5 parents as the breeder is unable to direct how the genomes combine or how they will interact with the environmental conditions. This unpredictability results in the expenditure of large amounts of research resources in the development of a superior new maize inbred line.

## 10 **Summary of the Invention**

According to the invention, there is provided a novel inbred maize line, designated PH48V. This invention thus relates to the seeds of inbred maize line PH48V, to the plants of inbred maize line PH48V, to methods for producing a maize  
15 plant produced by crossing the inbred maize line PH48V with itself or another maize line, and to methods for producing a maize plant containing in its genetic material one or more transgenes and to the transgenic maize plants produced by that method. This invention also relates to inbred maize lines derived from inbred maize line PH48V, to methods for producing other inbred maize lines derived from inbred maize line PH48V and to the inbred maize lines derived by the use of those methods. This invention  
20 further relates to hybrid maize seeds and plants produced by crossing the inbred line PH48V with another maize line.

## **Definitions**

In the description and examples that follow, a number of terms are used herein.  
25 In order to provide a clear and consistent understanding of the specification and claims, including the scope to be given such terms, the following definitions are provided. NOTE: ABS is in absolute terms and %MN is percent of the mean for the experiments in which the inbred or hybrid was grown. These designators will follow the descriptors to denote how the values are to be interpreted. Below are the descriptors used in the data  
30 tables included herein.

ANT ROT = ANTHRACNOSE STALK ROT (*Colletotrichum graminicola*). A 1 to 9 visual rating indicating the resistance to Anthracnose Stalk Rot. A higher score indicates a higher resistance.





EAR SZ = EAR SIZE. A 1 to 9 visual rating of ear size. The higher the rating the larger the ear size.

ECB 1LF = EUROPEAN CORN BORER FIRST GENERATION LEAF FEEDING (*Ostrinia nubilalis*). A 1 to 9 visual rating indicating the resistance to preflowering leaf feeding by first generation European Corn Borer. A higher score indicates a higher resistance.

ECB 2IT = EUROPEAN CORN BORER SECOND GENERATION INCHES OF TUNNELING (*Ostrinia nubilalis*). Average inches of tunneling per plant in the stalk.

ECB 2SC = EUROPEAN CORN BORER SECOND GENERATION (*Ostrinia nubilalis*). A 1 to 9 visual rating indicating post flowering degree of stalk breakage and other evidence of feeding by European Corn Borer, Second Generation. A higher score indicates a higher resistance.

ECB DPE = EUROPEAN CORN BORER DROPPED EARS (*Ostrinia nubilalis*). Dropped ears due to European Corn Borer. Percentage of plants that did not drop ears under second generation corn borer infestation.

EGR WTH = EARLY GROWTH. This is a measure of the relative height and size of a corn seedling at the 2-4 leaf stage of growth. This is a visual rating (1 to 9), with 1 being weak or slow growth, 5 being average growth and 9 being strong growth. Taller plants, wider leaves, more green mass and darker color constitute higher score.

EST CNT = EARLY STAND COUNT. This is a measure of the stand establishment in the spring and represents the number of plants that emerge on per plot basis for the inbred or hybrid.

EYE SPT = Eye Spot (*Kabatiella zeae* or *Aureobasidium zeae*). A 1 to 9 visual rating indicating the resistance to Eye Spot. A higher score indicates a higher resistance.

FUS ERS = FUSARIUM EAR ROT SCORE (*Fusarium moniliforme* or *Fusarium subglutinans*). A 1 to 9 visual rating indicating the resistance to Fusarium ear rot. A higher score indicates a higher resistance.

GDU = Growing Degree Units. Using the Barger Heat Unit Theory, which assumes that maize growth occurs in the temperature range 50°F - 86°F and that temperatures outside this range slow down growth; the maximum daily heat unit accumulation is 36 and the minimum daily heat unit accumulation is 0. The seasonal accumulation of GDU is a major factor in determining maturity zones.

GDU SHD = GDU TO SHED. The number of growing degree units (GDUs) or heat units required for an inbred line or hybrid to have approximately 50 percent of the plants shedding pollen and is measured from the time of planting. Growing degree units are calculated by the Barger Method, where the heat units for a 24-hour period are:

5 
$$\text{GDU} = \frac{(\text{Max. temp.} + \text{Min. temp.})}{2} - 50$$

The highest maximum temperature used is 86° F. and the lowest minimum temperature used is 50° F. For each inbred or hybrid it takes a certain number of GDUs to reach various stages of plant development.

GDU SLK = GDU TO SILK. The number of growing degree units required for an inbred line or hybrid to have approximately 50 percent of the plants with silk emergence from time of planting. Growing degree units are calculated by the Barger Method as given in GDU SHD definition.

15 GIB ERS = GIBBERELLA EAR ROT (PINK MOLD) (*Gibberella zeae*). A 1 to 9 visual rating indicating the resistance to Gibberella Ear Rot. A higher score indicates a higher resistance.

20 GLF SPT = Gray Leaf Spot (*Cercospora zeae-maydis*). A 1 to 9 visual rating indicating the resistance to Gray Leaf Spot. A higher score indicates a higher resistance.

GOS WLT = Goss' Wilt (*Corynebacterium nebraskense*). A 1 to 9 visual rating indicating the resistance to Goss' Wilt. A higher score indicates a higher resistance.

25 GRN APP = GRAIN APPEARANCE. This is a 1 to 9 rating for the general appearance of the shelled grain as it is harvested based on such factors as the color of harvested grain, any mold on the grain, and any cracked grain. High scores indicate good grain quality.

30 H/POP = YIELD AT HIGH DENSITY. Yield ability at relatively high plant densities on 1-9 relative rating system with a higher number indicating the hybrid responds well to high plant densities for yield relative to other hybrids. A 1, 5, and 9 would represent very poor, average, and very good yield response, respectively, to increased plant density.

HC BLT = HELMINTHOSPORIUM CARBONUM LEAF BLIGHT (*Helminthosporium carbonum*). A 1 to 9 visual rating indicating the resistance to Helminthosporium infection. A higher score indicates a higher resistance.

HD SMT = HEAD SMUT (*Sphacelotheca reiliana*). This score indicates the percentage of plants not infected.

HSK CVR = HUSK COVER. A 1 to 9 score based on performance relative to key checks, with a score of 1 indicating very short husks, tip of ear and kernels showing; 5 is intermediate coverage of the ear under most conditions, sometimes with thin husk; and a 9 has husks extending and closed beyond the tip of the ear. Scoring can best be done near physiological maturity stage or any time during dry down until harvested.

INC D/A = GROSS INCOME (DOLLARS PER ACRE). Relative income per acre assuming drying costs of two cents per point above 15.5 percent harvest moisture and current market price per bushel.

INCOME/ACRE. Income advantage of hybrid to be patented over other hybrid on per acre basis.

INC ADV = GROSS INCOME ADVANTAGE. GROSS INCOME advantage of variety #1 over variety #2.

KSZ DCD = KERNEL SIZE DISCARD. The percent of discard seed; calculated as the sum of discarded tip kernels and extra large kernels.

L/POP = YIELD AT LOW DENSITY. Yield ability at relatively low plant densities on a 1-9 relative system with a higher number indicating the hybrid responds well to low plant densities for yield relative to other hybrids. A 1, 5, and 9 would represent very poor, average, and very good yield response, respectively, to low plant density.

MDM CPX = MAIZE DWARF MOSAIC COMPLEX (MDMV = Maize Dwarf Mosaic Virus and MCDV = Maize Chlorotic Dwarf Virus). A 1 to 9 visual rating indicating the resistance to Maize Dwarf Mosaic Complex. A higher score indicates a higher resistance.

MST = HARVEST MOISTURE. The moisture is the actual percentage moisture of the grain at harvest.

MST ADV = MOISTURE ADVANTAGE. The moisture advantage of variety #1 over variety #2 as calculated by: MOISTURE of variety #2 - MOISTURE of variety #1 = MOISTURE ADVANTAGE of variety #1.

NLF BLT = Northern Leaf Blight (*Helminthosporium turcicum* or *Exserohilum turcicum*). A 1 to 9 visual rating indicating the resistance to Northern Leaf Blight. A higher score indicates a higher resistance.

OILT = GRAIN OIL. Absolute value of oil content of the kernel as predicted by Near-Infrared Transmittance and expressed as a percent of dry matter.

PLT HT = PLANT HEIGHT. This is a measure of the height of the plant from the ground to the tip of the tassel in inches.

POL SC = POLLEN SCORE. A 1 to 9 visual rating indicating the amount of pollen shed. The higher the score the more pollen shed.

5 POL WT = POLLEN WEIGHT. This is calculated by dry weight of tassels collected as shedding commences minus dry weight from similar tassels harvested after shedding is complete.

POP K/A = PLANT POPULATIONS. Measured as 1000s per acre.

10 POP ADV = PLANT POPULATION ADVANTAGE. The plant population advantage of variety #1 over variety #2 as calculated by PLANT POPULATION of variety #2 - PLANT POPULATION of variety #1 = PLANT POPULATION ADVANTAGE of variety #1.

15 PRM = PREDICTED RELATIVE MATURITY. This trait, predicted relative maturity, is based on the harvest moisture of the grain. The relative maturity rating is based on a known set of checks and utilizes standard linear regression analyses and is also referred to as the Comparative Relative Maturity Rating System that is similar to the Minnesota Relative Maturity Rating System.

20 PRM SHD = A relative measure of the growing degree units (GDU) required to reach 50% pollen shed. Relative values are predicted values from the linear regression of observed GDU's on relative maturity of commercial checks.

PROT = GRAIN PROTEIN. Absolute value of protein content of the kernel as predicted by Near-Infrared Transmittance and expressed as a percent of dry matter.

25 RT LDG = ROOT LODGING. Root lodging is the percentage of plants that do not root lodge; plants that lean from the vertical axis at an approximately 30° angle or greater would be counted as root lodged.

RTL ADV = ROOT LODGING ADVANTAGE. The root lodging advantage of variety #1 over variety #2.

30 SCT GRN = SCATTER GRAIN. A 1 to 9 visual rating indicating the amount of scatter grain (lack of pollination or kernel abortion) on the ear. The higher the score the less scatter grain.

SDG VGR = SEEDLING VIGOR. This is the visual rating (1 to 9) of the amount of vegetative growth after emergence at the seedling stage (approximately five leaves). A higher score indicates better vigor.

SEL IND = SELECTION INDEX. The selection index gives a single measure of the hybrid's worth based on information for up to five traits. A maize breeder may utilize his or her own set of traits for the selection index. One of the traits that is almost always included is yield. The selection index data presented in the tables represent the mean value averaged across testing stations.

SLF BLT = SOUTHERN LEAF BLIGHT (*Helminthosporium maydis* or *Bipolaris maydis*). A 1 to 9 visual rating indicating the resistance to Southern Leaf Blight. A higher score indicates a higher resistance.

SOU RST = SOUTHERN RUST (*Puccinia polysora*). A 1 to 9 visual rating indicating the resistance to Southern Rust. A higher score indicates a higher resistance.

STA GRN = STAY GREEN. Stay green is the measure of plant health near the time of black layer formation (physiological maturity). A high score indicates better late-season plant health.

STD ADV = STALK STANDING ADVANTAGE. The advantage of variety #1 over variety #2 for the trait STK CNT.

STK CNT = NUMBER OF PLANTS. This is the final stand or number of plants per plot.

STK LDG = STALK LODGING. This is the percentage of plants that did not stalk lodge (stalk breakage) as measured by either natural lodging or pushing the stalks and determining the percentage of plants that break below the ear.

STRT = GRAIN STARCH. Absolute value of starch content of the kernel as predicted by Near-Infrared Transmittance and expressed as a percent of dry matter.

STW WLT = Stewart's Wilt (*Erwinia stewartii*). A 1 to 9 visual rating indicating the resistance to Stewart's Wilt. A higher score indicates a higher resistance.

TAS BLS = TASSEL BLAST. A 1 to 9 visual rating was used to measure the degree of blasting (necrosis due to heat stress) of the tassel at the time of flowering. A 1 would indicate a very high level of blasting at time of flowering, while a 9 would have no tassel blasting.

TAS SZ = TASSEL SIZE. A 1 to 9 visual rating was used to indicate the relative size of the tassel. The higher the rating the larger the tassel.

TAS WT = TASSEL WEIGHT. This is the average weight of a tassel (grams) just prior to pollen shed.



TEX EAR = EAR TEXTURE. A 1 to 9 visual rating was used to indicate the relative hardness (smoothness of crown) of mature grain. A 1 would be very soft (extreme dent) while a 9 would be very hard (flinty or very smooth crown).

TILLER = TILLERS. A count of the number of tillers per plot that could possibly shed pollen was taken. Data are given as a percentage of tillers: number of tillers per plot divided by number of plants per plot.

TST WT = TEST WEIGHT (UNADJUSTED). The measure of the weight of the grain in pounds for a given volume (bushel).

TST WTA = TEST WEIGHT ADJUSTED. The measure of the weight of the grain in pounds for a given volume (bushel) adjusted for 15.5 percent moisture.

TSW ADV = TEST WEIGHT ADVANTAGE. The test weight advantage of variety #1 over variety #2.

WIN M% = PERCENT MOISTURE WINS.

WIN Y% = PERCENT YIELD WINS.

YLD = YIELD. It is the same as BU ACR ABS.

YLD ADV = YIELD ADVANTAGE. The yield advantage of variety #1 over variety #2 as calculated by: YIELD of variety #1 - YIELD variety #2 = yield advantage of variety #1.

YLD SC = YIELD SCORE. A 1 to 9 visual rating was used to give a relative rating for yield based on plot ear piles. The higher the rating the greater visual yield appearance.

#### **Detailed Description of the Invention**

Inbred maize lines are typically developed for use in the production of hybrid maize lines. Inbred maize lines need to be highly homogeneous, homozygous and reproducible to be useful as parents of commercial hybrids. There are many analytical methods available to determine the homozygotic and phenotypic stability of these inbred lines.

The oldest and most traditional method of analysis is the observation of phenotypic traits. The data is usually collected in field experiments over the life of the maize plants to be examined. Phenotypic characteristics most often observed are for traits associated with plant morphology, ear and kernel morphology, insect and disease resistance, maturity, and yield.

In addition to phenotypic observations, the genotype of a plant can also be examined. There are many laboratory-based techniques available for the analysis, comparison and characterization of plant genotype; among these are Isozyme Electrophoresis, Restriction Fragment Length Polymorphisms (RFLPs), Randomly Amplified Polymorphic DNAs (RAPDs), Arbitrarily Primed Polymerase Chain Reaction (AP-PCR), DNA Amplification Fingerprinting (DAF), Sequence Characterized Amplified Regions (SCARs), Amplified Fragment Length Polymorphisms (AFLPs), and Simple Sequence Repeats (SSRs) which are also referred to as Microsatellites.

The most widely used of these laboratory techniques are Isozyme Electrophoresis and RFLPs as discussed in Lee, M., "Inbred Lines of Maize and Their Molecular Markers," *The Maize Handbook*, (Springer-Verlag, New York, Inc. 1994, at 423- 432) incorporated herein by reference. Isozyme Electrophoresis is a useful tool in determining genetic composition, although it has relatively low number of available markers and the low number of allelic variants among maize inbreds. RFLPs have the advantage of revealing an exceptionally high degree of allelic variation in maize and the number of available markers is almost limitless.

Maize RFLP linkage maps have been rapidly constructed and widely implemented in genetic studies. One such study is described in Boppenmaier, et al., "Comparisons among strains of inbreds for RFLPs", *Maize Genetics Cooperative Newsletter*, 65:1991, pg. 90, is incorporated herein by reference. This study used 101 RFLP markers to analyze the patterns of 2 to 3 different deposits each of five different inbred lines. The inbred lines had been selfed from 9 to 12 times before being adopted into 2 to 3 different breeding programs. It was results from these 2 to 3 different breeding programs that supplied the different deposits for analysis. These five lines were maintained in the separate breeding programs by selfing or sibbing and rogueing off-type plants for an additional one to eight generations. After the RFLP analysis was completed, it was determined the five lines showed 0-2% residual heterozygosity. Although this was a relatively small study, it can be seen using RFLPs that the lines had been highly homozygous prior to the separate strain maintenance.

Inbred maize line PH48V is a yellow, dent maize inbred that is suited as a female for producing first generation F1 maize hybrids. Inbred maize line PH48V is best adapted to the Southeast region of the United States and can be used to produce hybrids from approximately 121 relative maturity based on the Comparative Relative Maturity Rating System for harvest moisture of grain. Inbred maize line PH48V

demonstrates high resistance to Southern Leaf Blight, high resistance to Northern Leaf Blight and Gray Leaf Spot as an inbred per se. In hybrid combination, including for its area of adaptation, inbred PH48V demonstrates high yield, above average resistance to stalk lodging, above average resistance to root lodging, strong staygreen, tall plant, high ear placement and strong disease resistance.

The inbred has shown uniformity and stability within the limits of environmental influence for all the traits as described in the Variety Description Information (Table 1) that follows. The inbred has been self-pollinated and ear-rowed a sufficient number of generations with careful attention paid to uniformity of plant type to ensure the homozygosity and phenotypic stability necessary to use in commercial production. The line has been increased both by hand and in isolated fields with continued observation for uniformity. No variant traits have been observed or are expected in PH48V.

Inbred maize line PH48V, being substantially homozygous, can be reproduced by planting seeds of the line, growing the resulting maize plants under self-pollinating or sib-pollinating conditions with adequate isolation, and harvesting the resulting seed, using techniques familiar to the agricultural arts.

TABLE 1  
VARIETY DESCRIPTION INFORMATION  
VARIETY = PH48V

1. TYPE: (describe intermediate types in Comments section):						
5	2	1=Sweet 2=Dent 3=Flint 4=Flour 5=Pop 6=Ornamental				
2. MATURITY:						
DAYS    HEAT UNITS						
	078	1,534.7	From emergence to 50% of plants in silk			
	076	1,486.7	From emergence to 50% of plants in pollen			
10	002	0,058.3	From 10% to 90% pollen shed			
			From 50% silk to harvest at 25% moisture			
3. PLANT:						
				Standard	Sample	
				Deviation	Size	
	0,226.7	cm	Plant Height (to tassel tip)	8.14	15	
15	0,086.0	cm	Ear Height (to base of top ear node)	7.55	15	
	0,014.3	cm	Length of Top Ear Internode	1.81	15	
	0.0		Average Number of Tillers	0.01	3	
	1.1		Average Number of Ears per Stalk	0.18	3	
	1.0		Anthocyanin of Brace Roots: 1=Absent 2=Faint 3=Moderate 4=Dark			
4. LEAF:						
				Standard	Sample	
				Deviation	Size	
	010.1	cm	Width of Ear Node Leaf	0.42	15	
	071.6	cm	Length of Ear Node Leaf	1.93	15	
	06.1		Number of leaves above top ear	0.12	15	
25	016.6		Degrees Leaf Angle (measure from 2nd leaf above ear at anthesis to stalk above leaf)	3.02	15	
	03		Leaf Color            Dark Green            (Munsell code)		5GY34	
	1.0		Leaf Sheath Pubescence (Rate on scale from 1=none to 9=like peach fuzz)			
			Marginal Waves (Rate on scale from 1=none to 9=many)			
30			Longitudinal Creases (Rate on scale from 1=none to 9=many)			
5. TASSEL:						
				Standard	Sample	
				Deviation	Size	
	09.9		Number of Primary Lateral Branches	0.31	15	
	034.7		Branch Angle from Central Spike	3.25	15	
35	51.5	cm	Tassel Length (from top leaf collar to tassel tip)	1.70	15	
	5.0		Pollen Shed (rate on scale from 0=male sterile to 9=heavy shed)			
	01		Anther Color            Light Green            (Munsell code)		2.5GY88	
	01		Glume Color            Light Green            (Munsell code)		7.5GY56	
	1.0		Bar Glumes (Glume Bands): 1=Absent 2=Present			
40	22	cm	Peduncle Length (cm. from top leaf to basal branches)			
6a. EAR (Unhusked Data):						
	1	Silk Color (3 days after emergence)		Light Green	(Munsell code)	2.5GY96
	1	Fresh Husk Color (25 days after 50% silking)		Light Green	(Munsell code)	5GY78
	21	Dry Husk Color (65 days after 50% silking)		Buff	(Munsell code)	5Y92
45	1	Position of Ear at Dry Husk Stage: 1= Upright 2= Horizontal 3= Pendant				
	8	Husk Tightness (Rate of Scale from 1=very loose to 9=very tight)				
	3	Husk Extension (at harvest): 1=Short (ears exposed) 2=Medium (<8 cm)				
		3=Long (8-10 cm beyond ear tip) 4=Very Long (>10 cm)				







### Further Embodiments of the Invention

This invention also is directed to methods for producing a maize plant by crossing a first parent maize plant with a second parent maize plant wherein either the first or second parent maize plant is an inbred maize plant of the line PH48V. Further, both first and second parent maize plants can come from the inbred maize line PH48V. Still further, this invention also is directed to methods for producing an inbred maize line PH48V-derived maize plant by crossing inbred maize line PH48V with a second maize plant and growing the progeny seed, and repeating the crossing and growing steps with the inbred maize line PH48V-derived plant from 0 to 5 times. Thus, any such methods using the inbred maize line PH48V are part of this invention: selfing, backcrosses, hybrid production, crosses to populations, and the like. All plants produced using inbred maize line PH48V as a parent are within the scope of this invention, including plants derived from inbred maize line PH48V. Advantageously, the inbred maize line is used in crosses with other, different, maize inbreds to produce first generation (F<sub>1</sub>) maize hybrid seeds and plants with superior characteristics.

A further embodiment of the invention is a single gene conversion or introgression of the maize plant disclosed herein in which the gene or genes of interest (encoding the desired trait) are introduced through traditional (non-transformation) breeding techniques, such as backcrossing (Hallauer *et al*, 1988). One or more genes may be introduced using these techniques. Desired traits transferred through this process include, but are not limited to, waxy starch, nutritional enhancements, industrial enhancements, disease resistance, insect resistance, herbicide resistance and yield enhancements. The gene of interest is transferred from the donor parent to the recurrent parent, in this case, the maize plant disclosed herein. These single gene traits may result from either the transfer of a dominant allele or a recessive allele. Selection of progeny containing the trait of interest is done by direct selection for a trait associated with a dominant allele. Selection of progeny for a trait that is transferred via a recessive allele, such as the waxy starch characteristic, requires growing and selfing the first backcross to determine which plants carry the recessive alleles. Recessive traits may require additional progeny testing in successive backcross generations to determine the presence of the gene of interest.

It should be understood that the inbred can, through routine manipulation of cytoplasmic or other factors, be produced in a male-sterile form. Such embodiments are also contemplated within the scope of the present claims.

As used herein, the term plant includes plant cells, plant protoplasts, plant cell tissue cultures from which maize plants can be regenerated, plant calli, plant clumps, and plant cells that are intact in plants or parts of plants, such as embryos, pollen, ovules, seeds, flowers, kernels, ears, cobs, leaves, husks, stalks, roots, root tips, anthers, silk and the like.

Duncan, Williams, Zehr, and Widholm, *Planta* (1985) 165:322-332 reflects that 97% of the plants cultured that produced callus were capable of plant regeneration. Subsequent experiments with both inbreds and hybrids produced 91% regenerable callus that produced plants. In a further study in 1988, Songstad, Duncan & Widholm in *Plant Cell Reports* (1988), 7:262-265 reports several media additions that enhance regenerability of callus of two inbred lines. Other published reports also indicated that "nontraditional" tissues are capable of producing somatic embryogenesis and plant regeneration. K.P. Rao, et al., *Maize Genetics Cooperation Newsletter*, 60:64-65 (1986), refers to somatic embryogenesis from glume callus cultures and B.V. Conger, et al., *Plant Cell Reports*, 6:345-347 (1987) indicates somatic embryogenesis from the tissue cultures of maize leaf segments. Thus, it is clear from the literature that the state of the art is such that these methods of obtaining plants are, and were, "conventional" in the sense that they are routinely used and have a very high rate of success.

Tissue culture of maize is described in European Patent Application, publication 160,390, incorporated herein by reference. Maize tissue culture procedures are also described in Green and Rhodes, "Plant Regeneration in Tissue Culture of Maize," *Maize for Biological Research* (Plant Molecular Biology Association, Charlottesville, Virginia 1982, at 367-372) and in Duncan, et al., "The Production of Callus Capable of Plant Regeneration from Immature Embryos of Numerous *Zea Mays* Genotypes," 165 *Planta* 322-332 (1985). Thus, another aspect of this invention is to provide cells which upon growth and differentiation produce maize plants having the physiological and morphological characteristics of inbred line PH48V.

The utility of inbred maize line PH0KT also extends to crosses with other species. Commonly, suitable species will be of the family Gramineae, and especially of the genera *Zea*, *Tripsacum*, *Coix*, *Schlerachne*, *Polytoca*, *Chionachne*, and *Trilobachne*, of the tribe Maydeae. Potentially suitable for crosses with PH48V may be the various varieties of grain sorghum, *Sorghum bicolor* (L.) Moench.

#### Transformation of Maize

With the advent of molecular biological techniques that have allowed the



resistance to the antibiotic hygromycin. Vanden Elzen *et al.*, *Plant Mol. Biol.*, 5: 299 (1985).

Additional selectable marker genes of bacterial origin that confer resistance to antibiotics include gentamycin acetyl transferase, streptomycin phosphotransferase, aminoglycoside- 3' -adenyl transferase, the bleomycin resistance determinant. Hayford *et al.*, *Plant Physiol.* 86: 1216 (1988), Jones *et al.*, *Mol. Gen. Genet.*, 210: 86 (1987), Svab *et al.*, *Plant Mol. Biol.*, 14: 197 (1990), Hille *et al.*, *Plant Mol. Biol.* 7: 171 (1986). Other selectable marker genes confer resistance to herbicides such as glyphosate, glufosinate or broxynil. Comai *et al.*, *Nature* 317: 741-744 (1985), Gordon-Kamm *et al.*, *Plant Cell* 2: 603-618 (1990) and Stalker *et al.*, *Science* 242: 419-423 (1988).

Other selectable marker genes for plant transformation are not of bacterial origin. These genes include, for example, mouse dihydrofolate reductase, plant 5 - enolpyruvylshikimate-3 -phosphate synthase and plant acetolactate synthase. Eichholtz *et al.*, *Somatic Cell Mol. Genet.* 13: 67 (1987), Shah *et al.*, *Science* 233: 478 (1986), Charest *et al.*, *Plant Cell Rep.* 8: 643 (1990).

Another class of marker genes for plant transformation require screening of presumptively transformed plant cells rather than direct genetic selection of transformed cells for resistance to a toxic substance such as an antibiotic. These genes are particularly useful to quantify or visualize the spatial pattern of expression of a gene in specific tissues and are frequently referred to as reporter genes because they can be fused to a gene or gene regulatory sequence for the investigation of gene expression. Commonly used genes for screening presumptively transformed cells include  $\beta$ -glucuronidase (*GUS*),  $\beta$ -galactosidase, luciferase and chloramphenicol acetyltransferase. Jefferson, R.A., *Plant Mol. Biol. Rep.* 5: 387 (1987), Teeri *et al.*, *EMBO J.* 8: 343 (1989), Koncz *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 84:131 (1987), De Block *et al.*, *EMBO J.* 3: 1681 (1984). Another approach to the identification of relatively rare transformation events has been use of a gene that encodes a dominant constitutive regulator of the *Zea mays* anthocyanin pigmentation pathway. Ludwig *et al.*, *Science* 247: 449 (1990).

Recently, *in vivo* methods for visualizing GUS activity that do not require destruction of plant tissue have been made available. Molecular Probes Publication 2908, Imagene Green™, p. 1-4 (1993) and Naleway *et al.*, *J. Cell Biol.* 115: 151a (1991). However, these *in vivo* methods for visualizing GUS activity have not proven useful for recovery of transformed cells because of low sensitivity, high fluorescent backgrounds,



and limitations associated with the use of luciferase genes as selectable markers.

More recently, a gene encoding Green Fluorescent Protein (GFP) has been utilized as a marker for gene expression in prokaryotic and eukaryotic cells. Chalfie *et al.*, *Science* 263: 802 (1994). GFP and mutants of GFP may be used as screenable markers.

### Promoters

Genes included in expression vectors must be driven by a nucleotide sequence comprising a regulatory element, for example, a promoter. Several types of promoters are now well known in the transformation arts, as are other regulatory elements that can be used alone or in combination with promoters.

As used herein "promoter" includes reference to a region of DNA upstream from the start of transcription and involved in recognition and binding of RNA polymerase and other proteins to initiate transcription. A "plant promoter" is a promoter capable of initiating transcription in plant cells. Examples of promoters under developmental control include promoters that preferentially initiate transcription in certain tissues, such as leaves, roots, seeds, fibers, xylem vessels, tracheids, or sclerenchyma. Such promoters are referred to as "tissue-preferred". Promoters which initiate transcription only in certain tissues are referred to as "tissue-specific". A "cell type" specific promoter primarily drives expression in certain cell types in one or more organs, for example, vascular cells in roots or leaves. An "inducible" promoter is a promoter which is under environmental control. Examples of environmental conditions that may effect transcription by inducible promoters include anaerobic conditions or the presence of light. Tissue-specific, tissue-preferred, cell type specific, and inducible promoters constitute the class of "non-constitutive" promoters. A "constitutive" promoter is a promoter which is active under most environmental conditions.

#### A. Inducible Promoters

An inducible promoter is operably linked to a gene for expression in maize. Optionally, the inducible promoter is operably linked to a nucleotide sequence encoding a signal sequence which is operably linked to a gene for expression in maize. With an inducible promoter the rate of transcription increases in response to an inducing agent.

Any inducible promoter can be used in the instant invention. See Ward *et al.* *Plant Mol. Biol.* 22: 361-366 (1993). Exemplary inducible promoters include, but are not limited to, that from the ACEI system which responds to copper (Mett *et al.* *PNAS* 90: 4567-4571 (1993)); In2 gene from maize which responds to benzenesulfonamide

herbicide safeners (Hershey *et al.*, *Mol. Gen. Genetics* 227: 229-237 (1991) and Gatz *et al.*, *Mol. Gen. Genetics* 243: 32-38 (1994)) or Tet repressor from Tn10 (Gatz *et al.*, *Mol. Gen. Genet.* 227: 229-237 (1991). A particularly preferred inducible promoter is a promoter that responds to an inducing agent to which plants do not normally respond.

5 An exemplary inducible promoter is the inducible promoter from a steroid hormone gene, the transcriptional activity of which is induced by a glucocorticosteroid hormone. Schena *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 88: 0421 (1991).

#### **B. Constitutive Promoters**

A constitutive promoter is operably linked to a gene for expression in maize or

10 the constitutive promoter is operably linked to a nucleotide sequence encoding a signal sequence which is operably linked to a gene for expression in maize.

Many different constitutive promoters can be utilized in the instant invention. Exemplary constitutive promoters include, but are not limited to, the promoters from plant viruses such as the 35S promoter from CaMV (Odell *et al.*, *Nature* 313: 810-812

15 (1985) and the promoters from such genes as rice actin (McElroy *et al.*, *Plant Cell* 2: 163-171 (1990)); ubiquitin (Christensen *et al.*, *Plant Mol. Biol.* 12: 619-632 (1989) and Christensen *et al.*, *Plant Mol. Biol.* 18: 675-689 (1992)); pEMU (Last *et al.*, *Theor. Appl. Genet.* 31: 581-588 (1991)); MAS (Velten *et al.*, *EMBO J.* 3: 2723-2730 (1984)) and maize H3 histone (Lepetit *et al.*, *Mol. Gen. Genet.* 231: 276-285 (1992) and Atanassova

20 *et al.*, *Plant Journal* 2 (3): 291-300 (1992)).

The ALS promoter, a XbaI/NcoI fragment 5' to the *Brassica napus* ALS3 structural gene (or a nucleotide sequence that has substantial sequence similarity to said XbaI/NcoI fragment), represents a particularly useful constitutive promoter. See PCT application WO96/30530.

#### **C. Tissue-specific or Tissue-Preferred Promoters**

A tissue-specific promoter is operably linked to a gene for expression in maize. Optionally, the tissue-specific promoter is operably linked to a nucleotide sequence encoding a signal sequence which is operably linked to a gene for expression in maize. Plants transformed with a gene of interest operably linked to a tissue-specific promoter

30 produce the protein product of the transgene exclusively, or preferentially, in a specific tissue.

Any tissue-specific or tissue-preferred promoter can be utilized in the instant invention. Exemplary tissue-specific or tissue-preferred promoters include, but are not limited to, a root-preferred promoter, such as that from the phaseolin gene (Murai *et al.*,

Science 23: 476-482 (1983) and Sengupta-Gopalan *et al.*, *Proc. Natl. Acad. Sci. USA* 82: 3320-3324 (1985)); a leaf-specific and light-induced promoter such as that from *cab* or *rubisco* (Simpson *et al.*, *EMBO J.* 4(11): 2723-2729 (1985) and Timko *et al.*, *Nature* 318: 579-582 (1985)); an anther-specific promoter such as that from *LAT52* (Twell *et al.*, *Mol. Gen. Genet.* 217: 240-245 (1989)); a pollen-specific promoter such as that from *Zm13* (Guerrero *et al.*, *Mol. Gen. Genet.* 224: 161-168 (1993)) or a microspore-preferred promoter such as that from *apg* (Twell *et al.*, *Sex. Plant Reprod.* 6: 217-224 (1993)).

### Signal Sequences For Targeting Proteins to Subcellular Compartments

Transport of protein produced by transgenes to a subcellular compartment such as the chloroplast, vacuole, peroxisome, glyoxysome, cell wall or mitochondrion, or for secretion into the apoplast, is accomplished by means of operably linking the nucleotide sequence encoding a signal sequence to the 5' and/or 3' region of a gene encoding the protein of interest. Targeting sequences at the 5' and/or 3' end of the structural gene may determine, during protein synthesis and processing, where the encoded protein is ultimately compartmentalized. The presence of a signal sequence directs a polypeptide to either an intracellular organelle or subcellular compartment or for secretion to the apoplast. Many signal sequences are known in the art. See, for example, Becker *et al.*, *Plant Mol. Biol.* 20: 49 (1992), Close, P.S., Master's Thesis, Iowa State University (1993), Knox, C., *et al.*, "Structure and Organization of Two Divergent Alpha-Amylase Genes From Barley", *Plant Mol. Biol.* 9: 3-17 (1987), Lerner *et al.*, *Plant Physiol.* 91: 124-129 (1989), Fontes *et al.*, *Plant Cell* 3: 483-496 (1991), Matsuoka *et al.*, *Proc. Natl. Acad. Sci.* 88: 834 (1991), Gould *et al.*, *J. Cell Biol* 108: 1657 (1989), Creissen *et al.*, *Plant J.* 2: 129 (1991), Kalderon, D., Robers, B., Richardson, W., and Smith A., "A short amino acid sequence able to specify nuclear location", *Cell* 39: 499-509 (1984), Stiefel, V., Ruiz-Avila, L., Raz R., Valles M., Gomez J., Pages M., Martinez-Izquierdo J., Ludevid M., Landale J., Nelson T., and Puigdomenech P., "Expression of a maize cell wall hydroxyproline-rich glycoprotein gene in early leaf and root vascular differentiation", *Plant Cell* 2: 785-793 (1990).

### Foreign Protein Genes and Agronomic Genes

With transgenic plants according to the present invention, a foreign protein can be produced in commercial quantities. Thus, techniques for the selection and propagation of transformed plants, which are well understood in the art, yield a plurality of transgenic plants which are harvested in a conventional manner, and a foreign protein then can be extracted from a tissue of interest or from total biomass. Protein



DNA molecules encoding  $\delta$ -endotoxin genes can be purchased from American Type Culture Collection (Rockville, MD), for example, under ATCC Accession Nos. 40098, 67136, 31995 and 31998.

(C) A lectin. See, for example, the disclosure by Van Damme *et al.*, *Plant Molec. Biol.* 24: 25 (1994), who disclose the nucleotide sequences of several *Clivia miniata* mannose-binding lectin genes.

(D) A vitamin-binding protein such as avidin. See PCT application US93/06487 the contents of which are hereby incorporated by. The application teaches the use of avidin and avidin homologues as larvicides against insect pests.

(E) An enzyme inhibitor, for example, a protease inhibitor or an amylase inhibitor. See, for example, Abe *et al.*, *J. Biol. Chem.* 262: 16793 (1987) (nucleotide sequence of rice cysteine proteinase inhibitor), Huub *et al.*, *Plant Molec. Biol.* 21: 985 (1993) (nucleotide sequence of cDNA encoding tobacco proteinase inhibitor I), and Sumitani *et al.*, *Biosci. Biotech. Biochem.* 57: 1243 (1993) (nucleotide sequence of *Streptomyces nitrosporeus*  $\alpha$ -amylase inhibitor).

(F) An insect-specific hormone or pheromone such as an ecdysteroid and juvenile hormone, a variant thereof, a mimetic based thereon, or an antagonist or agonist thereof. See, for example, the disclosure by Hammock *et al.*, *Nature* 344: 458 (1990), of baculovirus expression of cloned juvenile hormone esterase, an inactivator of juvenile hormone.

(G) An insect-specific peptide or neuropeptide which, upon expression, disrupts the physiology of the affected pest. For example, see the disclosures of Regan, *J. Biol. Chem.* 269: 9 (1994) (expression cloning yields DNA coding for insect diuretic hormone receptor), and Pratt *et al.*, *Biochem. Biophys. Res. Comm.* 163: 1243 (1989) (an allostatin is identified in *Diploptera punctata*). See also U.S. patent No.5,266,317 to Tomalski *et al.*, who disclose genes encoding insect-specific, paralytic neurotoxins.

(H) An insect-specific venom produced in nature by a snake, a wasp, *etc.* For example, see Pang *et al.*, *Gene* 116: 165 (1992), for disclosure of heterologous expression in plants of a gene coding for a scorpion insectotoxic peptide.

(I) An enzyme responsible for an hyperaccumulation of a monerpene, a sesquiterpene, a steroid, hydroxamic acid, a phenylpropanoid derivative or another non-protein molecule with insecticidal activity.

(J) An enzyme involved in the modification, including the post-translational



modification, of a biologically active molecule; for example, a glycolytic enzyme, a proteolytic enzyme, a lipolytic enzyme, a nuclease, a cyclase, a transaminase, an esterase, a hydrolase, a phosphatase, a kinase, a phosphorylase, a polymerase, an elastase, a chitinase and a glucanase, whether natural or synthetic. See PCT application WO 93/02197 in the name of Scott *et al.*, which discloses the nucleotide sequence of a callase gene. DNA molecules which contain chitinase-encoding sequences can be obtained, for example, from the ATCC under Accession Nos. 39637 and 67152. See also Kramer *et al.*, *Insect Biochem. Molec. Biol.* 23: 691 (1993), who teach the nucleotide sequence of a cDNA encoding tobacco hookworm chitinase, and Kawalleck *et al.*, *Plant Molec. Biol.* 21: 673 (1993), who provide the nucleotide sequence of the parsley *ubi4-2* polyubiquitin gene.

(K) A molecule that stimulates signal transduction. For example, see the disclosure by Botella *et al.*, *Plant Molec. Biol.* 24: 757 (1994), of nucleotide sequences for mung bean calmodulin cDNA clones, and Griess *et al.*, *Plant Physiol.* 104: 1467 (1994), who provide the nucleotide sequence of a maize calmodulin cDNA clone.

(L) A hydrophobic moment peptide. See PCT application WO95/16776 (disclosure of peptide derivatives of Tachyplesin which inhibit fungal plant pathogens) and PCT application WO95/18855 (teaches synthetic antimicrobial peptides that confer disease resistance), the respective contents of which are hereby incorporated by reference.

(M) A membrane permease, a channel former or a channel blocker. For example, see the disclosure by Jaynes *et al.*, *Plant Sci.* 89: 43 (1993), of heterologous expression of a cecropin- $\beta$  lytic peptide analog to render transgenic tobacco plants resistant to *Pseudomonas solanacearum*.

(N) A viral-invasive protein or a complex toxin derived therefrom. For example, the accumulation of viral coat proteins in transformed plant cells imparts resistance to viral infection and/or disease development effected by the virus from which the coat protein gene is derived, as well as by related viruses. See Beachy *et al.*, *Ann. Rev. Phytopathol.* 28: 451 (1990). Coat protein-mediated resistance has been conferred upon transformed plants against alfalfa mosaic virus, cucumber mosaic virus, tobacco streak virus, potato virus X, potato virus Y, tobacco etch virus, tobacco rattle virus and tobacco mosaic virus. *Id.*

(O) An insect-specific antibody or an immunotoxin derived therefrom. Thus, an antibody targeted to a critical metabolic function in the insect gut would inactivate an

affected enzyme, killing the insect. Cf. Taylor *et al.*, Abstract #497, SEVENTH INT'L SYMPOSIUM ON MOLECULAR PLANT-MICROBE INTERACTIONS (Edinburgh, Scotland, 1994) (enzymatic inactivation in transgenic tobacco via production of single-chain antibody fragments).

5 (P) A virus-specific antibody. See, for example, Tavladoraki *et al.*, *Nature* 366: 469 (1993), who show that transgenic plants expressing recombinant antibody genes are protected from virus attack.

(Q) A developmental-arrestive protein produced in nature by a pathogen or a parasite. Thus, fungal endo  $\alpha$ -1,4-D-polygalacturonases facilitate fungal colonization  
10 and plant nutrient release by solubilizing plant cell wall homo- $\alpha$ -1,4-D-galacturonase. See Lamb *et al.*, *Bio/Technology* 10: 1436 (1992). The cloning and characterization of a gene which encodes a bean endopolygalacturonase-inhibiting protein is described by Toubart *et al.*, *Plant J.* 2: 367 (1992).

(R) A developmental-arrestive protein produced in nature by a plant. For  
15 example, Logemann *et al.*, *Bio/Technology* 10: 305 (1992), have shown that transgenic plants expressing the barley ribosome-inactivating gene have an increased resistance to fungal disease.

## 2. Genes That Confer Resistance To A Herbicide, For Example:

(A) A herbicide that inhibits the growing point or meristem, such as an  
20 imidazalinone or a sulfonylurea. Exemplary genes in this category code for mutant ALS and AHAS enzyme as described, for example, by Lee *et al.*, *EMBO J.* 7: 1241 (1988), and Miki *et al.*, *Theor. Appl. Genet.* 80: 449 (1990), respectively.

(B) Glyphosate (resistance imparted by mutant 5-enolpyruvyl-3-phosphokimate  
synthase (EPSP) and *aroA* genes, respectively) and other phosphono compounds  
25 such as glufosinate (phosphinothricin acetyl transferase (PAT) and *Streptomyces hygroscopicus* phosphinothricin acetyl transferase (*bar*) genes), and pyridinoxy or phenoxy proprionic acids and cyclohexones (ACCase inhibitor-encoding genes). See, for example, U.S. patent No. 4,940,835 to Shah *et al.*, which discloses the nucleotide sequence of a form of EPSP which can confer glyphosate resistance. A DNA molecule  
30 encoding a mutant *aroA* gene can be obtained under ATCC accession No. 39256, and the nucleotide sequence of the mutant gene is disclosed in U.S. patent No. 4,769,061 to Comai. European patent application No. 0 333 033 to Kumada *et al.* and U.S. patent No. 4,975,374 to Goodman *et al.* disclose nucleotide sequences of glutamine synthetase genes which confer resistance to herbicides such as L-phosphinothricin.



*licheniformis*  $\alpha$ -amylase), Elliot *et al.*, *Plant Molec. Biol.* 21: 515 (1993) (nucleotide sequences of tomato invertase genes), Sogaard *et al.*, *J. Biol. Chem.* 268: 22480 (1993) (site-directed mutagenesis of barley  $\alpha$ -amylase gene), and Fisher *et al.*, *Plant Physiol.* 102: 1045 (1993) (maize endosperm starch branching enzyme II).

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## Methods for Maize Transformation

Numerous methods for plant transformation have been developed, including biological and physical, plant transformation protocols. See, for example, Miki *et al.*, "Procedures for Introducing Foreign DNA into Plants" in *Methods in Plant Molecular Biology and Biotechnology*, Glick, B.R. and Thompson, J.E. Eds. (CRC Press, Inc., Boca Raton, 1993) pages 67-88. In addition, expression vectors and *in vitro* culture methods for plant cell or tissue transformation and regeneration of plants are available. See, for example, Gruber *et al.*, "Vectors for Plant Transformation" in *Methods in Plant Molecular Biology and Biotechnology*, Glick, B.R. and Thompson, J.E. Eds. (CRC Press, Inc., Boca Raton, 1993) pages 89-119.

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### A. *Agrobacterium*-mediated Transformation

One method for introducing an expression vector into plants is based on the natural transformation system of *Agrobacterium*. See, for example, Horsch *et al.*, *Science* 227: 1229 (1985). *A. tumefaciens* and *A. rhizogenes* are plant pathogenic soil bacteria which genetically transform plant cells. The Ti and Ri plasmids of *A. tumefaciens* and *A. rhizogenes*, respectively, carry genes responsible for genetic transformation of the plant. See, for example, Kado, C.I., *Crit. Rev. Plant. Sci.* 10: 1 (1991). Descriptions of *Agrobacterium* vector systems and methods for *Agrobacterium*-mediated gene transfer are provided by Gruber *et al.*, *supra*, Miki *et al.*, *supra*, and Moloney *et al.*, *Plant Cell Reports* 8: 238 (1989). See also, U.S. Patent No. 5,591,616, issued Jan. 7, 1997.

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### B. Direct Gene Transfer

Despite the fact the host range for *Agrobacterium*-mediated transformation is broad, some major cereal crop species and gymnosperms have generally been recalcitrant to this mode of gene transfer, even though some success has recently been achieved in rice and maize. Hiei *et al.*, *The Plant Journal* 6: 271-282 (1994); U.S. Patent No. 5,591,616, issued Jan. 7, 1997. Several methods of plant transformation, collectively referred to as direct gene transfer, have been developed as an alternative to *Agrobacterium*-mediated transformation.

30





herein, "crossing" can refer to a simple X by Y cross, or the process of backcrossing, depending on the context.

### **Industrial Applicability**

Maize is used as human food, livestock feed, and as raw material in industry.  
 5 The food uses of maize, in addition to human consumption of maize kernels, include both products of dry- and wet-milling industries. The principal products of maize dry milling are grits, meal and flour. The maize wet-milling industry can provide maize starch, maize syrups, and dextrose for food use. Maize oil is recovered from maize germ, which is a by-product of both dry- and wet-milling industries.

10 Maize, including both grain and non-grain portions of the plant, is also used extensively as livestock feed, primarily for beef cattle, dairy cattle, hogs, and poultry.

Industrial uses of maize include production of ethanol, maize starch in the wet-milling industry and maize flour in the dry-milling industry. The industrial applications of maize starch and flour are based on functional properties, such as viscosity, film  
 15 formation, adhesive properties, and ability to suspend particles. The maize starch and flour have application in the paper and textile industries. Other industrial uses include applications in adhesives, building materials, foundry binders, laundry starches, explosives, oil-well muds, and other mining applications.

Plant parts other than the grain of maize are also used in industry: for example,  
 20 stalks and husks are made into paper and wallboard and cobs are used for fuel and to make charcoal.

The seed of inbred maize line PH48V, the plant produced from the inbred seed, the hybrid maize plant produced from the crossing of the inbred, hybrid seed, and various parts of the hybrid maize plant and transgenic versions of the foregoing, can be  
 25 utilized for human food, livestock feed, and as a raw material in industry.

### **Performance Examples of PH48V**

In the examples that follow, the traits and characteristics of inbred maize line PH48V are given as a line. The data collected on inbred maize line PH48V is presented  
 30 for the key characteristics and traits.



[illegible]

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[illegible]

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inbred PHKV1 crossed to PHN46. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid presents higher than average ear placement and a

significantly taller plant than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid shows significantly better staygreen scores than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging and brittle stalk than the PHKV1/PHN46 hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PHKV1/PHN46 hybrid.

The results in Table 4B compare inbred PH48V crossed to inbred PH0KT and inbred PHW52 crossed to PHK46. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields with significantly higher test weight of grain than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores and demonstrates above average resistance to root lodging. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging and brittle stalk than the PHW52/PHK46 hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PHW52/PHK46 hybrid.

The results in Table 4C compare inbred PH48V crossed to inbred PH0KT and inbred PH6M1 crossed to PH24M. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to stalk lodging than the PH6M1/PH24M hybrid. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PH6M1/PH24M hybrid and above average resistance to Northern Leaf Blight.

The results in Table 4D compare inbred PH48V crossed to inbred PH0KT and inbred PH07D crossed to PH0V0. The results show the PH48V/PH0KT hybrid to demonstrate above average and significantly higher yields than the PH07D/PH0V0 hybrid. The PH48V/PH0KT hybrid presents a significantly taller plant and a significantly higher ear placement than the PH07D/PH0V0 hybrid. The PH48V/PH0KT hybrid shows above average staygreen scores. The PH48V/PH0KT hybrid demonstrates above average and significantly better resistance to brittle stalk than the PH07D/PH0V0 hybrid,

and shows above average resistance to stalk lodging. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Southern Leaf Blight than the PH07D/PH0V0 hybrid and above average resistance to Gray Leaf Spot and Northern Leaf Blight.

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- 10       above average staygreen scores and demonstrates above average resistance to stalk lodging. The PH48V/PH0KT hybrid shows above average and significantly better resistance to Gray Leaf Spot and Southern Leaf Blight than the PH07D/PH24M hybrid and above average resistance to Northern Leaf Blight.

TABLE 2A  
PAIRED INBRED COMPARISON REPORT

VARIETY #1 = PH48V  
VARIETY #2 = PHOOM

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		BU ACR ABS	BU ACR %MN	MST ABS	TST WT ABS	EGR WTH ABS	EST CNT ABS	TIL LER ABS	GDU SHD ABS	GDU SLK ABS
TOTAL SUM	1	71.2	105	26.1	56.0	4.8	38.5	2.4	148.7	154.6
	2	57.6	91	26.1	54.5	4.4	35.5	1.3	147.7	154.8
	LOCS	12	12	14	2	9	22	13	20	20
	REPS	13	13	15	2	10	23	13	20	20
	DIFF	13.5	14	0.0	1.5	0.4	3.0	1.1	1.0	0.3
	PR > T	.111	.403	.999	.066*	.211	.018+	.276	.347	.838

		POL SC ABS	TAS SZ ABS	PLT HT ABS	EAR HT ABS	RT LDG ABS	STA GRN ABS	STK LDG ABS	BRT STK ABS	SCT GRN ABS
TOTAL SUM	1	5.7	5.3	92.7	39.5	98.7	7.3	100.0	100.0	7.0
	2	4.7	4.6	85.7	34.0	100.0	6.5	93.1	100.0	3.0
	LOCS	3	13	6	4	3	4	2	1	1
	REPS	3	13	6	4	3	4	2	1	1
	DIFF	1.0	0.7	7.0	5.5	1.3	0.8	6.9	0.0	4.0
	PR > T	.000#	.056*	.001#	.086*	.423	.486	.500		

		EAR SZ ABS	TEX EAR ABS	EAR MLD ABS	BAR PLT ABS	GLF SPT ABS	STW WLT ABS	COM RST ABS	CLD TST ABS	CLD TST %MN
TOTAL SUM	1	5.0	6.0	7.0	98.4	9.0	8.0	5.3	78.0	108
	2	4.0	5.5	4.5	92.2	8.5	8.0	5.5	63.7	87
	LOCS	1	2	2	11	2	1	4	3	3
	REPS	1	2	2	11	2	1	4	3	3
	DIFF	1.0	0.5	2.5	6.2	0.5	0.0	0.3	14.3	21
	PR > T		.500	.344	.060*	.500		.391	.246	.258

		KSZ DCD ABS
TOTAL SUM	1	5.7
	2	5.3
	LOCS	3
	REPS	3
	DIFF	0.3
	PR > T	.945

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

TABLE 2B  
PAIRED INBRED COMPARISON REPORT

VARIETY #1 = PH48V  
VARIETY #2 = PHKV1

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		BU ACR ABS	BU ACR %MN	MST ABS	TST WT ABS	EGR WTH ABS	EST CNT ABS	TIL LER ABS	GDU SHD ABS	GDU SLK ABS
TOTAL SUM	1	72.5	109	26.6	54.8	5.0	30.9	2.7	151.5	157.6
	2	64.5	96	23.7	55.1	4.8	31.2	1.5	155.8	157.5
	LOCS	35	35	39	9	26	47	40	56	54
	REPS	37	37	41	10	27	48	41	56	54
	DIFF	8.0	13	2.9	0.3	0.1	0.3	1.2	4.2	0.1
	PR > T	.066*	.061*	.000#	.472	.622	.594	.068*	.000#	.883

		POL SC ABS	TAS SZ ABS	PLT HT ABS	EAR HT ABS	RT LDG ABS	STA GRN ABS	STK LDG ABS	BRT STK ABS	SCT GRN ABS
TOTAL SUM	1	5.7	5.4	95.7	41.2	96.3	7.8	100.0	100.0	6.8
	2	4.3	2.7	89.2	38.7	89.7	4.7	80.8	100.0	6.4
	LOCS	3	36	25	13	5	11	2	3	5
	REPS	3	36	25	13	5	12	2	3	5
	DIFF	1.3	2.8	6.5	2.5	6.5	3.1	19.2	0.0	0.4
	PR > T	.057*	.000#	.000#	.018+	.257	.001#	.500	.999	.374

		EAR SZ ABS	TEX EAR ABS	EAR MLD ABS	BAR PLT ABS	GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS
TOTAL SUM	1	5.0	6.0	7.6	94.5	7.9	6.5	6.0	7.3	4.5
	2	6.0	6.0	7.6	89.7	2.6	3.5	7.0	5.5	4.5
	LOCS	1	2	5	36	9	1	1	3	2
	REPS	1	2	5	36	12	2	2	4	3
	DIFF	1.0	0.0	0.0	4.9	5.4	3.0	1.0	1.8	0.0
	PR > T		.999	.999	.062*	.000#			.008#	.999

		HD SMT ABS	MDM CPX ABS	FUS ERS ABS	DIP ERS ABS	COM RST ABS	CLD TST ABS	CLD TST %MN	KSZ DCD ABS
TOTAL SUM	1	100.0	4.0	6.7	1.5	5.3	85.2	107	3.6
	2	100.0	4.0	7.2	2.5	1.3	83.3	105	1.8
	LOCS	1	1	6	1	4	15	15	15
	REPS	2	2	7	2	4	15	15	15
	DIFF	0.0	0.0	0.5	1.0	4.0	1.9	2	1.8
	PR > T			.296		.011+	.670	.751	.039+

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

TABLE 2C  
PAIRED INBRED COMPARISON REPORT

VARIETY #1 = PH48V  
VARIETY #2 = PH07D

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		BU	BU		TST	EGR	EST	TIL	GDU	GDU
		ACR	ACR	MST	WT	WTH	CNT	LER	SHD	SLK
		ABS	%MN	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL	SUM	1	73.9	108	26.4	55.1	4.9	28.1	2.9	151.4
		2	65.9	98	20.6	57.1	5.7	27.9	2.5	149.1
	LOCS	33	33	37	8	31	51	44	66	66
	REPS	35	35	39	9	32	65	45	66	68
	DIFF	8.0	10	5.8	2.0	0.8	0.3	0.3	2.4	6.4
	PR > T	.067*	.129	.000#	.009#	.005#	.645	.613	.000#	.000#

		POL	POL	POL	TAS	TAS	PLT	EAR	RT	STA
		WT	WT	SC	BLS	SZ	HT	HT	LDG	GRN
		ABS	%MN	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL	SUM	1	178.3	83	5.7	3.0	5.6	95.7	40.9	96.3
		2	186.7	87	4.7	1.0	4.0	89.6	37.9	98.5
	LOCS	2	2	3	1	42	33	15	5	13
	REPS	2	2	3	1	42	33	15	5	14
	DIFF	8.3	4	1.0	2.0	1.5	6.1	3.0	2.2	1.9
	PR > T	.430	.427	.000#		.000#	.000#	.031+	.204	.000#

		STK	BRT	SCT	EAR	TEX	EAR	BAR	GLF	NLF
		LDG	STK	GRN	SZ	EAR	MLD	PLT	SPT	BLT
		ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
TOTAL	SUM	1	100.0	100.0	6.4	5.0	6.0	7.3	94.7	7.9
		2	78.8	96.6	7.3	5.0	6.5	6.9	92.9	6.6
	LOCS	2	5	8	2	3	8	41	12	3
	REPS	2	8	8	2	3	8	41	17	5
	DIFF	21.2	3.4	0.9	0.0	0.5	0.4	1.7	1.3	1.3
	PR > T	.500	.220	.021+	.999	.225	.442	.315	.000#	.270

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG



[illegible]

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TABLE 3

Average Inbred By Tester Performance Comparing PH48V To PH07D Crossed  
To The Same Inbred Testers And Grown In The Same Experiments.

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		BU ACR ABS	BU ACR %MN	MST %MN	EGR WTH %MN	EST CNT %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN
TOTAL SUM	REPS	21	21	21	5	8	20	9	9	4	10
	LOCS	21	21	21	5	8	20	9	9	4	10
	PH48V	170	114	102	85	105	104	101	110	103	107
	PH07D	156	105	93	93	97	100	99	100	103	127
	DIFF	14	9	9	7	8	3	2	9	0	20
	PR > T	0.00	0.00	0.00	0.50	0.01	0.08	0.34	0.01	0.99	0.09

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		STK LDG %MN	SLF BLT ABS	COM RST ABS	SOU RST ABS	DRP EAR %MN	GLF SPT ABS
TOTAL SUM	REPS	13	1	3	1	1	1
	LOCS	13	1	3	1	1	1
	PH48V	104	7	7	4	100	7
	PH07D	105	6	6	6	100	5
	DIFF	1	1	1	2	0	2
	PR > T	0.56		0.23		0.99	

\*PR > T values are valid only for comparisons with Locs >= 10.

TABLE 4A  
INBREDS IN HYBRID COMBINATION REPORT

		VARIETY #1 = PH48V/PH0KT VARIETY #2 = PHKV1/PHN46								
		PRM	PRM	BU	BU		TST	EGR	EST	GDU
		ABS	SHD	ACR	ACR	MST	WT	WTH	CNT	SHD
		ABS	ABS	ABS	%MN	%MN	ABS	%MN	%MN	%MN
10	TOTAL SUM	1	119	116	173.7	106	105	57.0	92	103
		2	115	117	166.9	101	94	57.2	92	104
15	LOCS		7	6	143	143	58	27	30	26
	REPS		7	6	165	165	59	30	34	29
	DIFF		4	1	6.8	4	11	0.3	0	1
	PR > T		.002#	.007#	.001#	.001#	.000#	.252	.999	.733
		GDU	STK	PLT	EAR	RT	STA	STK	BRT	DRP
		SLK	CNT	HT	HT	LDG	GRN	LDG	STK	EAR
		%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN	%MN
25	TOTAL SUM	1	101	101	104	108	99	109	102	108
		2	102	102	100	107	103	86	97	98
30	LOCS		17	190	51	50	31	47	55	14
	REPS		20	245	62	60	34	52	59	19
	DIFF		2	1	4	2	4	23	6	11
	PR > T		.014+	.183	.000#	.195	.027+	.000#	.002#	.037+
		GLF	NLF	SLF	STW	ANT	HD		MDM	FUS
		SPT	BLT	BLT	WLT	ROT	SMT	CLN	CPX	ERS
		ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS	ABS
40	TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0
		2	2.7	5.8	3.6	5.8	2.9	79.6	6.8	3.0
45	LOCS		8	5	7	5	6	2	2	1
	REPS		12	7	10	5	10	4	4	2
	DIFF		3.9	0.0	3.8	1.2	2.1	17.3	0.5	0.0
	PR > T		.000#	.999	.000#	.235	.021+	.500	.500	.548
		* = 10% SIG + = 5% SIG # = 1% SIG								

TABLE 4A (cont.)  
INBREDS IN HYBRID COMBINATION REPORT

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VARIETY #1 = PH48V/PHOKT  
VARIETY #2 = PHKV1/PHN46

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\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

TABLE 4B  
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PHOKT  
VARIETY #2 = PHW52/PHK46

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		PRM ABS	PRM SHD ABS	BU ACR ABS	BU ACR %MN	MST %MN	TST WT ABS	EGR WTH %MN	EST CNT %MN	GDU SHD %MN
TOTAL SUM	1	120	115	176.6	106	106	57.0	91	103	100
	2	118	116	167.1	100	102	56.0	99	100	101
	LOCS	5	5	125	125	125	66	25	33	23
	REPS	5	5	140	140	140	67	27	37	25
	DIFF	1	0	9.5	6	4	1.0	7	3	1
	PR > T	.006#	.999	.000#	.000#	.000#	.000#	.091*	.098*	.009#

		GDU SLK %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN	STK LDG %MN	BRT STK %MN	DRP EAR %MN
TOTAL SUM	1	100	101	104	108	101	111	103	108	100
	2	101	98	100	101	98	107	97	90	100
	LOCS	15	160	40	40	13	40	38	13	9
	REPS	17	197	42	42	13	43	40	17	11
	DIFF	1	2	4	7	3	4	5	18	0
	PR > T	.203	.007#	.000#	.000#	.384	.458	.010+	.023+	.999

		GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS	HD SMT ABS	CLN ABS	MDM CPX ABS	FUS ERS ABS
TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0	5.1
	2	4.4	5.3	6.5	4.2	4.1	72.9	6.3	3.0	6.2
	LOCS	8	4	7	5	6	2	2	1	9
	REPS	12	6	10	5	10	4	4	2	12
	DIFF	2.2	0.5	0.9	2.8	0.9	24.0	0.0	0.0	1.1
	PR > T	.000#	.423	.007#	.009#	.186	.426	.999		.159

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

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TABLE 4B (cont.)  
INBREDS IN HYBRID COMBINATION REPORT

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VARIETY #1 = PH48V/PHOKT  
VARIETY #2 = PHW52/PHK46

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\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG



TABLE 4C  
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PH0KT  
VARIETY #2 = PH6M1/PH24M

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		PRM ABS	PRM SHD ABS	BU ACR ABS	BU ACR %MN	MST %MN	TST WT ABS	EGR WTH %MN	EST CNT %MN	GDU SHD %MN
TOTAL SUM	1	119	115	176.8	104	106	57.4	95	104	100
	2	115	116	163.6	96	94	58.9	105	101	102
	LOCS	4	3	78	78	78	35	14	17	17
	REPS	4	3	93	93	93	36	16	21	19
	DIFF	4	1	13.2	8	11	1.5	10	3	2
	PR > T	.002#	.022+	.000#	.000#	.000#	.000#	.051*	.134	.001#

		GDU SLK %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN	STK LDG %MN	BRT STK %MN	DRP EAR %MN
TOTAL SUM	1	100	101	105	109	100	112	102	108	100
	2	103	100	102	100	102	96	98	100	100
	LOCS	12	114	29	28	7	25	22	13	8
	REPS	14	157	36	34	7	28	24	17	10
	DIFF	2	1	3	9	2	16	4	8	0
	PR > T	.003#	.337	.011+	.000#	.326	.064*	.004#	.278	.999

		GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS	HD SMT ABS		MDM CPX ABS	FUS ERS ABS
TOTAL SUM	1	6.5	5.8	7.6	7.0	5.0	96.9	6.3	3.0	5.3
	2	3.9	4.4	6.7	6.0	4.4	77.1	6.5	3.0	5.3
	LOCS	7	4	5	5	6	2	2	1	8
	REPS	11	6	8	5	10	4	4	2	10
	DIFF	2.6	1.4	0.9	1.0	0.6	19.8	0.3	0.0	0.0
	PR > T	.003#	.062*	.001#	.142	.363	.500	.500		.999

		DIP ERS ABS	COM RST ABS	HSK CVR ABS	HSK CVR %MN	OIL T ABS	PRO T ABS	STR T ABS
TOTAL SUM	1	2.8	7.3	5.1	93	4.4	9.0	72.3
	2	3.5	6.3	5.5	101	4.2	9.7	71.6
	LOCS	2	2	8	8	8	8	8
	REPS	4	3	8	8	8	8	8
	DIFF	0.8	1.0	0.4	8	0.2	0.6	0.8
	PR > T	.205	.000#	.351	.351	.215	.232	.004#

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

TABLE 4D  
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PH0KT  
VARIETY #2 = PH07D/PH0V0

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		PRM ABS	PRM SHD ABS	BU ACR ABS	BU ACR %MN	MST %MN	TST WT ABS	EGR WTH %MN	EST CNT %MN	GDU SHD %MN
TOTAL SUM	1	119	116	169.3	104	105	57.2	96	104	100
	2	118	116	155.1	95	103	57.5	104	102	101
	LOCS	5	4	94	94	94	34	18	17	20
	REPS	5	4	108	108	108	35	21	21	23
	DIFF	1	0	14.2	9	3	0.3	8	3	1
	PR > T	.327	.999	.000#	.000#	.002#	.224	.026+	.202	.203

		GDU SLK %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN	STK LDG %MN	BRT STK %MN	DRP EAR %MN
TOTAL SUM	1	101	101	105	108	98	110	102	108	100
	2	101	103	101	101	100	108	99	90	100
	LOCS	14	136	37	36	25	32	38	14	8
	REPS	17	182	48	46	28	37	42	19	10
	DIFF	0	1	3	7	2	2	3	18	0
	PR > T	.999	.085*	.001#	.000#	.296	.802	.177	.030+	.999

		GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS	HD SMT ABS	CLN ABS	MDM CPX ABS	FUS ERS ABS
TOTAL SUM	1	6.5	5.8	7.6	7.0	5.0	96.9	6.3	3.0	5.0
	2	5.9	6.4	5.9	6.4	4.7	100.0	6.5	3.0	6.4
	LOCS	7	5	5	5	6	2	2	1	9
	REPS	11	7	8	5	10	4	4	2	11
	DIFF	0.6	0.6	1.7	0.6	0.3	3.1	0.3	0.0	1.4
	PR > T	.122	.109	.010+	.208	.655	.500	.500		.069*

		DIP ERS ABS	COM RST ABS	HSK CVR ABS	HSK CVR %MN	OIL T ABS	PRO T ABS	STR T ABS
TOTAL SUM	1	3.3	6.1	5.1	93	4.3	9.4	72.1
	2	3.5	5.2	6.1	111	4.2	9.1	71.8
	LOCS	2	7	8	8	7	7	7
	REPS	3	8	8	8	7	7	7
	DIFF	0.3	0.9	1.0	19	0.1	0.4	0.3
	PR > T	.874	.045+	.033+	.034+	.687	.551	.293

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

TABLE 4E  
INBREDS IN HYBRID COMBINATION REPORT

VARIETY #1 = PH48V/PH0KT  
VARIETY #2 = PH07D/PH24M

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		PRM ABS	PRM SHD ABS	BU ACR ABS	BU ACR %MN	MST %MN	TST WT ABS	EGR WTH %MN	EST CNT %MN	GDU SHD %MN
TOTAL SUM	1	119	115	169.9	105	105	57.4	93	103	100
	2	114	115	160.8	100	94	59.3	103	102	101
	LOCS	6	4	119	119	120	49	21	29	20
	REPS	6	4	134	134	135	50	23	33	22
	DIFF	5	0	9.2	6	11	1.9	10	1	1
	PR > T	.000#	.999	.000#	.000#	.000#	.000#	.042+	.449	.161

		GDU SLK %MN	STK CNT %MN	PLT HT %MN	EAR HT %MN	RT LDG %MN	STA GRN %MN	STK LDG %MN	BRT STK %MN	DRP EAR %MN
TOTAL SUM	1	100	101	104	109	99	112	102	108	100
	2	101	102	100	96	102	102	104	104	100
	LOCS	12	159	42	41	25	35	49	13	9
	REPS	14	207	52	50	25	38	51	17	11
	DIFF	0	0	4	13	3	10	2	4	0
	PR > T	.999	.999	.000#	.000#	.089*	.097*	.042+	.406	.999

		GLF SPT ABS	NLF BLT ABS	SLF BLT ABS	STW WLT ABS	ANT ROT ABS	HD SMT ABS	CLN ABS	MDM CPX ABS	FUS ERS ABS
TOTAL SUM	1	6.6	5.8	7.4	7.0	5.0	96.9	6.3	3.0	4.8
	2	5.5	5.5	5.3	6.0	4.9	93.8	6.5	3.0	4.8
	LOCS	8	4	6	5	6	2	2	1	8
	REPS	12	6	8	5	10	4	4	2	11
	DIFF	1.1	0.3	2.1	1.0	0.1	3.1	0.3	0.0	0.1
	PR > T	.046+	.638	.000#	.230	.910	.500	.500		.945

		DIP ERS ABS	COM RST ABS	SOU RST ABS	HSK CVR ABS	HSK CVR %MN	OIL T ABS	PRO T ABS	STR T ABS
TOTAL SUM	1	3.3	6.2	4.0	5.1	92	4.3	9.3	71.8
	2	3.0	6.0	4.0	5.2	94	4.0	9.7	70.9
	LOCS	2	10	1	15	15	3	3	3
	REPS	3	10	1	15	15	3	3	3
	DIFF	0.3	0.2	0.0	0.1	2	0.2	0.4	0.9
	PR > T	.795	.443		.653	.673	.113	.414	.058*

\* = 10% SIG  
+ = 5% SIG  
# = 1% SIG

### Deposits

A deposit of the seed of inbred PH48V is and has been maintained by Pioneer Hi-Bred International, Inc., 800 Capital Square, 400 Locust Street, Des Moines, Iowa 50309-2340, since prior to the filing date of this application. Access to this deposit will be available during the pendency of the application to the Commissioner of Patents and Trademarks and person determined by the Commissioner to be entitled thereto upon request. Upon allowance of any claims in the application, the Applicant(s) will make available to the public without restriction a deposit of at least 2500 seeds of inbred PH48V with the American Type Culture Collection (ATCC), Manassas, Virginia, 20110. The seeds deposited with the ATCC will be taken from the same deposit maintained at Pioneer Hi-Bred and described above. Additionally, Applicant(s) will meet all the requirements of 37 C.F.R. §1.801 - 1.809, including providing an indication of the viability of the sample when the deposit is made. This deposit of Inbred Maize Line PH48V will be maintained in the ATCC Depository, which is a public depository, for a period of 30 years, or 5 years after the most recent request, or for the enforceable life of the patent, whichever is longer, and will be replaced if it ever becomes nonviable during that period. Applicant will impose no restrictions on the availability of the deposited material from the ATCC; however, Applicant has no authority to waive any restrictions imposed by law on the transfer of biological material or its transportation in commerce. Applicant does not waive any infringement of its rights granted under this patent or under the Plant Variety Protection Act (7 USC 2321 et seq.).

The foregoing invention has been described in detail by way of illustration and example for purposes of clarity and understanding. However, it will be obvious that certain changes and modifications such as single gene conversions and mutations, somoclonal variants, variant individuals selected from large populations of the plants of the instant inbred and the like may be practiced within the scope of the invention, as limited only by the scope of the appended claims.